

SEQUENCE LISTING

<110> Merck & Co., Inc.
Freedman, Leonard P.
Glantschnig, Helmut
Harada, Shun-ichi
Hess, John W.

<120> CYNOMOLGUS MONKEY DICKKOPF-4,
NUCLEOTIDES ENCODING SAME, AND USES THEREOF

<130> 21351Y

<150> 60/520,569

<151> 2003-11-17

<160> 13

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 675

<212> DNA

<213> MACACA FASCICULARIS

<400> 1

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tgcctgtctg acacagactg caataccaga aagttctgtcc tccagtccca caatgagaag 180
ccgttctgtg ctacatgtcg tgggttgagc aggaggtgcc agcgagatgc catgtgctgc 240
cctggggacac tctgcatgaa tgatgtttgt actacgatgg aagacgcaac cccaaaattg 300
gaaaggcagc ttgatgagca agatggcaca catgcagaag taacaactgg gcacccagtc 360
caggaaaacc aacccaagag gaagccaagt attaagaaat cacaaggcag gaagggacaa 420
gagggagaaaa gttgtctgag aacttttgac tgtggccctg gactttgctg tgctcgtcat 480
ttttggacga aaatttgtaa gccagtcctt ttggagggac aggtctgctc caggagaggg 540
cataaagaca ctgctcaagc tccagaaatc ttccagcgtt gcgactgtgg ccccgacta 600
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<210> 2

<211> 224

<212> PRT

<213> MACACA FASCICULARIS

<400> 2

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          20             25             30
Leu Gly Ala Arg Lys Gly Ser Gln Cys Leu Ser Asp Thr Asp Cys Asn
          35             40             45
Thr Arg Lys Phe Cys Leu Gln Ser His Asn Glu Lys Pro Phe Cys Ala
          50             55             60
Thr Cys Arg Gly Leu Gln Arg Arg Cys Gln Arg Asp Ala Met Cys Cys
65             70             75             80
Pro Gly Thr Leu Cys Met Asn Asp Val Cys Thr Thr Met Glu Asp Ala
          85             90             95

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Thr Pro Lys Leu Glu Arg Gln Leu Asp Glu Gln Asp Gly Thr His Ala
 100 105 110
 Glu Val Thr Thr Gly His Pro Val Gln Glu Asn Gln Pro Lys Arg Lys
 115 120 125
 Pro Ser Ile Lys Lys Ser Gln Gly Arg Lys Gly Gln Glu Gly Glu Ser
 130 135 140
 Cys Leu Arg Thr Phe Asp Cys Gly Pro Gly Leu Cys Cys Ala Arg His
 145 150 155 160
 Phe Trp Thr Lys Ile Cys Lys Pro Val Leu Leu Glu Gly Gln Val Cys
 165 170 175
 Ser Arg Arg Gly His Lys Asp Thr Ala Gln Ala Pro Glu Ile Phe Gln
 180 185 190
 Arg Cys Asp Cys Gly Pro Gly Leu Leu Cys Arg Ser Gln Leu Thr Ser
 195 200 205
 Asn Gln Gln His Ala Arg Leu Arg Val Cys Gln Lys Ile Glu Lys Leu
 210 215 220

<210> 3
 <211> 841
 <212> DNA
 <213> HOMO SAPIENS

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 gtggcggccg tccgtctggg gctgagctgg ctctgctctc ccctgggagc tctggctcctg 180
 gacttcaaca acatcaggag ctctgctgac ctgcatgggg cccggaaggg ctcacagtgc 240
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 gggacactct gtgtgaacga tgtttgtact acgatggaag atgcaacccc aatattagaa 420
 aggcagcttg atgagcaaga tggcacacat gcagaaggaa caactgggca cccagtccag 480
 gaaaaccaac caaaaggaa gccaaagtatt aagaaatcac aaggcaggaa gggacaagag 540
 ggagaaagtt gtctgagaac ttttgactgt ggccctggac tttgctgtgc tcgtcatttt 600
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<210> 4
 <211> 224
 <212> PRT
 <213> HOMO SAPIENS

<400> 4
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 His Gly Ala Arg Lys Gly Ser Gln Cys Leu Ser Asp Thr Asp Cys Asn
 35 40 45
 Thr Arg Lys Phe Cys Leu Gln Pro Arg Asp Glu Lys Pro Phe Cys Ala
 50 55 60
 Thr Cys Arg Gly Leu Arg Arg Arg Cys Gln Arg Asp Ala Met Cys Cys
 65 70 75 80
 Pro Gly Thr Leu Cys Val Asn Asp Val Cys Thr Thr Met Glu Asp Ala
 85 90 95

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Thr Pro Ile Leu Glu Arg Gln Leu Asp Glu Gln Asp Gly Thr His Ala
      100      105      110
Glu Gly Thr Thr Gly His Pro Val Gln Glu Asn Gln Pro Lys Arg Lys
      115      120      125
Pro Ser Ile Lys Lys Ser Gln Gly Arg Lys Gly Gln Glu Gly Glu Ser
      130      135      140
Cys Leu Arg Thr Phe Asp Cys Gly Pro Gly Leu Cys Cys Ala Arg His
      145      150      155      160
Phe Trp Thr Lys Ile Cys Lys Pro Val Leu Leu Glu Gly Gln Val Cys
      165      170      175
Ser Arg Arg Gly His Lys Asp Thr Ala Gln Ala Pro Glu Ile Phe Gln
      180      185      190
Arg Cys Asp Cys Gly Pro Gly Leu Cys Arg Ser Gln Leu Thr Ser
      195      200      205
Asn Arg Gln His Ala Arg Leu Arg Val Cys Gln Lys Ile Glu Lys Leu
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<210> 5
<211> 221
<212> PRT
<213> MUS MUSCULUS

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<400> 5
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      20      25      30
Gln Gly Ala Gly Lys Gly Ser Leu Cys Ala Ser Asp Arg Asp Cys Ser
      35      40      45
Glu Gly Lys Phe Cys Leu Ala Phe His Asp Glu Arg Ser Phe Cys Ala
      50      55      60
Thr Cys Arg Arg Val Arg Arg Arg Cys Gln Arg Ser Ala Val Cys Cys
      65      70      75      80
Pro Gly Thr Val Cys Val Asn Asp Val Cys Thr Ala Val Glu Asp Thr
      85      90      95
Arg Pro Val Met Asp Arg Asn Thr Asp Gly Gln Asp Gly Ala Tyr Ala
      100      105      110
Glu Gly Thr Thr Lys Trp Pro Ala Glu Glu Asn Arg Pro Gln Gly Lys
      115      120      125
Pro Ser Thr Lys Lys Ser Gln Ser Ser Lys Gly Gln Glu Gly Glu Ser
      130      135      140
Cys Leu Arg Thr Ser Asp Cys Gly Pro Gly Leu Cys Cys Ala Arg His
      145      150      155      160
Phe Trp Thr Lys Ile Cys Lys Pro Val Leu Arg Glu Gly Gln Val Cys
      165      170      175
Ser Arg Arg Gly His Lys Asp Thr Ala Gln Ala Pro Glu Ile Phe Gln
      180      185      190
Arg Cys Asp Cys Gly Pro Gly Leu Thr Cys Arg Ser Gln Val Thr Ser
      195      200      205
Asn Arg Gln His Ser Arg Leu Arg Val Cys Gln Arg Ile
      210      215      220

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<210> 6
<211> 24
<212> DNA
<213> Artificial Sequence

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<220>
<223> F2 PRIMER

<400> 6
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<210> 7
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> R2 PRIMER

<400> 7
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<210> 8
<211> 31
<212> DNA
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<220>
<223> F PRIMER

<400> 8
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<210> 9
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> FF PRIMER

<400> 9
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<210> 10
<211> 42
<212> DNA
<213> Artificial Sequence

<220>
<223> R PRIMER

<400> 10
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<210> 11
<211> 47
<212> DNA
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<223> RR PRIMER

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<210> 12
<211> 22
<212> DNA
<213> Artificial Sequence
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<223> Fseq PRIMER
<400> 12
gggacactct gcatgaatga tg 22
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<223> Rseq PRIMER
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